PubMed   Nucleotide   Protein   Genome   Structure   PopSet   Taxonomy   OMIM   Books

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Display   default   Save   Text   Add to Clipboard   Preview/Index   History   Clipboard   Details

ProbeSet, PubMed, Taxonomy, LinkOut

☐ 1: AA056177. zk70e10.r1 Soares...[gi:1548515]

IDENTIFIERS

dbEST Id:	681952
EST name:	zk70e10.r1
GenBank Acc:	AA056177
GenBank gi:	1548515
GDB Id:	3802789

CLONE INFO

Clone Id:	IMAGE:488202 (5')
Source:	IMAGE Consortium, LLNL
Insert length:	618
DNA type:	cDNA

PRIMERS

Sequencing:	-28M13 rev2 from Amersham
PolyA Tail:	Unknown

SEQUENCE

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TCAGGATGACCATAACTGGAGAAGNGAGTTGTATGCCCAAACAGTCTCTGAACGGTTACA
GGCCCATGGAAAGGAAAAACCCCAGATCATCATCTGTTACCCTGGGACTGGGCATTACATCGA
GCCTCCTTACTTCCCCCTGTGCCCAGCTTCCCTTCACAGATTACTGAACAAACATGTTAT
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AGCCTTCTTCTGCAAACACCTGGGAGGTACCCAGAAAACAGCTGTCCCTAAATTGTAATG
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GTGAATCAGATGTCTCCTGGATAACATTAAAGCCATGTCTTGT
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Quality: High quality sequence stops at base: 491

Entry Created: Sep 19 1996
Last Updated: May 11 1997

COMMENTS

This clone is available royalty-free through LLNL ; contact
the IMAGE Consortium (info@image.llnl.gov) for further
information.
Putative full length read
The vector to vector length is 584

PUTATIVE ID   Assigned by submitter
TR:G532505 G532505 BILE ACID COA: AMINO ACID
N-ACYLTRANSFERASE. ;

LIBRARY

Lib Name: Soares_pregnant_uterus_NbHPU
Organism: Homo sapiens
Sex: female
Organ: uterus
Develop. stage: adult
Lab host: DH10B
Vector: pT7T3-Pac
R. Site 1:   Not I
R. Site 2:   Eco RI
Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer
                 [5' AACTGGAAGAATTCGGCGGCCGCGCCTTTTTTTTTTTTTTTTTTT 3'],
                 double-stranded cDNA was ligated to Eco RI adaptors
                 (Pharmacia), digested with Not I and cloned into the Not I
                 and Eco RI sites of the modified pT7T3 vector. Library went
                 through one round of normalization. Library constructed by
                 M. Fatima Bonaldo.

SUBMITTER

Name: Wilson RK
Institution: Washington University School of Medicine
Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
E-mail: est@watson.wustl.edu

CITATIONS

Medline UID: 97044478
Title: Generation and analysis of 280,000 human expressed sequence
                 tags

Citation:

Authors: Hillier,L., Lennon,G., Becker,M., Bonaldo,M.F., Chiapelli,B.
, Chissoe,S., Dietrich,N., DuBuque,T., Favello,A., Gish,W.,
Hawkins,M., Hultman,M., Kucaba,T., Lacy,M., Le,M., Le,N.,
Mardis,E., Moore,B., Morris,M., Parsons,J., Prange,C.,
Rifkin,L., Rohlfing,T., Schellenberg,K., Soares,M.B., Tan,F.
, Thierry-Meg,J., Trevaskis,E., Underwood,K., Wohldmann,P.,
Waterston,R., Wilson,R., Marra,M.
Genome Res. 6 (9): 807-28 1996
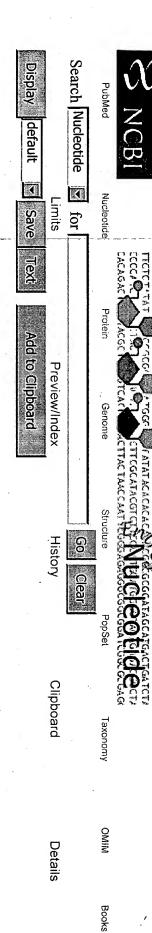
MAP DATA

Revised: October 24, 2001.

NCBI

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Nucleotide

Search [Nucleotide ▼] for [   ] Limits Preview/Index History Clipboard Details

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[Save] [Text] [Add to Clipboard]

[Display] [default ▼]

Taxonomy, LinkOut

☐ 1: AA639200. ns04b01.r1|NCI_CG...[gi:2562979]

IDENTIFIERS

dbEST Id:	**1342636**
EST name:	ns04b01.r1
GenBank Acc:	AA639200
GenBank gi:	2562979

CLONE INFO

Clone Id:	IMAGE:1182601
Source:	NCI
DNA type:	cDNA

PRIMERS

| Sequencing: | -28m13 rev1 ET from Amersham |
| PolyA Tail: | Unknown |

SEQUENCE

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CTACTACTACTAAATTCGCGGGATGTCGACCTCGTGGACATCGTGGATATAAGGAATGCT
CTCGTAGGAGGGTACAAGAGAACCCAGCACATGATGATTCAATAGAGAAGGCCCAGGCCAATCC
TGCTCATTGTTGGTCAGGATGACATAACTGGAGAAGTGAGTTGTATGCCCAAACAGTCTC
TGAACGGTTACAGGCCCATGGAAAGGAAAAACCCCAGATCATCGTTACCCTGGGACTGG
GCATTACATCGAGCCTCCTTACTTCCCCGTGCCCAGCTTCCCTTCACAGATTACTGAA
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GAAGCAAATTCTAGCCTTCTTCTGCAAACACCTGGGAGGTACCCAGAAAAACAGCTGTCC
TAAATTGTAATGCATTTGTCTTGTTGTTGACATGAGAGAATCAAGATCAGATTCTAGTGTT
CAGTAACCCTATGTGAATCAGATGTCTCCTGGATAACATTAAAGCCATGTCTTTGTCAT
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| Quality: | High quality sequence stops at base: 476 |

| Entry Created: | Oct 23 1997 |
| Last Updated: | Oct 23 1997 |

COMMENTS

Tissue Procurement: Lee Helman, M.D., Michael R. Emmert-Buck
, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing
Center
Clone distribution: NCI-CGAP clone distribution information
can be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html

PUTATIVE ID   Assigned by submitter
TR:G532505 G532505 BILE ACID COA: AMINO ACID
N-ACYLTRANSFERASE. ;

LIBRARY
Lib Name: NCI_CGAP_Ew1
Organism: Homo sapiens
Tissue type: Ewing's sarcoma
Lab host: DH10B
Vector: pAMP10
Description: mRNA made from Ewing's sarcoma, cDNA made by oligo-dT
priming. Non-directionally cloned. Size-selected on agarose
gel, average insert size 600 bp. Reference: Krizman et al.
(1996) Cancer Research 56:5380-5383.

SUBMITTER
Name: Robert Strausberg, Ph.D.
E-mail: cgapbs-r@mail.nih.gov

CITATIONS
Title: National Cancer Institute, Cancer Genome Anatomy Project
(CGAP), Tumor Gene Index
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
Year: 1997
Status: Unpublished

MAP DATA

Search [Nucleotide ▾] for [＿＿＿＿＿＿＿] [Go] [Clear]

PubMed   Nucleotide   Protein   Genome   Structure   PopSet   Taxonomy   OMIM   Books

[Display] [default ▾] [Save] [Text] [Add to Clipboard]   Preview/Index   History   Clipboard   Details   Taxonomy, LinkOut

☐ 1: AI830209. wj78f01.x1 NCI_CG...[gi:5450869]

IDENTIFIERS

dbEST Id: 2918151
EST name: wj78f01.x1
GenBank Acc: AI830209
GenBank gi: 5450869

CLONE INFO
Clone Id: IMAGE:2408953 (3')
Source: NCI
Insert length: 1242
DNA type: cDNA

PRIMERS
Sequencing: -40UP from Gibco
PolyA Tail: Unknown.

SEQUENCE

ATGACAAAGACACATGTCTTTAATGTTATCCAGGAGAGACATCTGATTCACATAGGGTTACTGA
ACACTAGAAATCTGATCTTGAATCTCTCAGTCAACAACAGACAAATGCATTACAATTTAG
GGACAGCGTGTTTTCTGGGTACCTCCCAGGTGTTTGCAGAAGAAGCTAGAGAATTTGCTTCC
AGGCATCTTCCTGGCCTTAGAATGAGCCCTCCACCCCATATAACATGTTTGT
TCAGTAAATCTGTGAAAGGAAGCTGGGCACAGGGGAAGTAAGGAGGCTCGATGTAATG
CCCAGTCCCAGGGTAACAGATGATCTGGGGTTTTTCCTTTCCATGGGCCTGTAACCGTTC
AGAGACTGTTTGGGCATAACAACTCACTTCTCCAGTTATGGTCATCCTGACCAACAATGAG
CAGGATGGGGCCCCTGGGCCTTCTCTATTGGAATCATGCTGGGGTTCTTGTACCCTCCTAC
GAGAGCAATTCCTTATATCCACAATGTCCACGAGGCCTGAGAAAGCTACCTTGATTCTCCT
CANGTCATAGCCCAATGGTGGAATGCTACTGTTGCTTATAGTTGATGGCTGTGTTCCACT
GATCCCAGATCCATTGATGGGAACTGTGGCTGAGACATTCTTNCAGAATGANGCCATTGA
GAGACANATATCAGCTCCTAGAGAAATGCCCAAATGCCTGGGCCTTTTAC

Quality: High quality sequence stops at base: 457

Entry Created: Jul 12 1999
Last Updated: Dec 21 1999

COMMENTS

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing
Center
Clone distribution: NCI-CGAP clone distribution information
can be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html

PUTATIVE ID   Assigned by submitter
               SW:YZ28_HUMAN P49753 HYPOTHETICAL PROTEIN ZAP128 ;

LIBRARY

Lib Name: NCI_CGAP_Lu19
Organism: Homo sapiens
Organ: lung
Tissue type: squamous cell carcinoma, poorly differentiated (4 pooled
                tumors, including primary and metastatic)
Develop. stage: adult
Lab host: DH10B (phage-resistant)
Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker
Description: 1st strand cDNA was prepared from pooled lung tumor tissue,
                and was then primed with a Not I - oligo(dT) primer.
                Double-stranded cDNA was ligated to Eco RI adaptors
                (Pharmacia), digested with Not I and cloned into the Not I
                and Eco RI sites of the modified pT7T3 vector. Library went
                through one round of normalization. Library constructed by
                Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D.
E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project
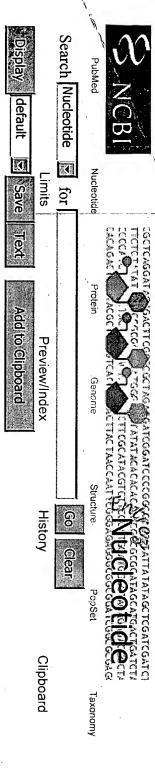   (CGAP), Tumor Gene Index
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
Year: 1997

NCBI Sequence Viewer

Status: Unpublished

MAP DATA

Revised: October 24, 2001.